RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	<u>/o/538,530</u>
Source:	IFW1.6
Date Processed by STIC:	2/22/06

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IFW16

RAW SEQUENCE LISTING DATE: 02/22/2006
PATENT APPLICATION: US/10/538,530 TIME: 12:11:07

Input Set : A:\MBP-017xxSeqList.txt
Output Set: N:\CRF4\02222006\J538530.raw

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3 <110> APPLICANT: Reski, Ralf
         Decker, Eva
         Justine Kiessling
 7 <120> TITLE OF INVENTION: FtsZ-POLYPEPTIDES AS A TARGET FOR HERBICIDAL COMPOUNDS
 9 <130> FILE REFERENCE: MBP-017XX
11 <140> CURRENT APPLICATION NUMBER: US 10/538,530
12 <141> CURRENT FILING DATE: 2005-06-10
14 <150> PRIOR APPLICATION NUMBER: PCT/EP2003/014162
15 <151> PRIOR FILING DATE: 2003-12-12
17 <150> PRIOR APPLICATION NUMBER: US 60/433,556
18 <151> PRIOR FILING DATE: 2002-12-13
20 <150> PRIOR APPLICATION NUMBER: US 60/438,466
21 <151> PRIOR FILING DATE: 2003-01-07
23 <160> NUMBER OF SEO ID NOS: 5
25 <170> SOFTWARE: PatentIn version 3.1
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28 <211> LENGTH: 1335
29 <212> TYPE: DNA
30 <213> ORGANISM: Physcomitrella patens
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37 tgtgggtgct tgagagcggg gaataagctg gataaggacc aatttgtggg tgatgggaaa
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39 ccacttatgc atcaacagac gcggggatgg agtcaggggc gggagaggtg tcacgcaggg
41 aggtctgtgg tgatggccag tatgagtggc gccaagatca aggtcattgg tgtaggcggc
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43 gggggcaaca atgctgtgaa ccgcatgatt gggagcggca ttcagggtgt tgatttttgg
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45 gccatcaaca cagatgttca agctttgcag aaatcacaag ccgaacatcg cgttcaaatc
                                                                         420
47 ggcgaagett tgacccgagg acttggtact ggtggaaage catteettgg agaacaagea
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49 gcagaggaat cgatagaaat cattgcacag gcagtggtag atgctgatct tgtcttcatt
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51 actgegggea tgggtggtgg aacggggtct ggggctgccc cggtcgttgc ccgtgtggcc
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53 aaagaggcag ggcaactcac tgttggtgtt gtcacttatc cgtttacgtt tgagggccgt
                                                                         660
55 cggagaagcc agcaggcagt ggaggcaata gagaatctgc ggaagtctgt cgacagtctt
                                                                         720
57 attgtcattc ctaatgaccg tctactcgat gtctccggag ataaaactcc tcttcaggaa
                                                                         780
59 gcattttctc tagccgacga tgttcttagg cagggagttc aaggcatttc agacatcatc
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61 acaacgccag gtcttgtgaa tgttgatttt gcagatgtta gagctgtaat gagtaactca
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63 ggtacagcca tgcttggcgt tggctcctct agtggcaaga atcgtgctga ggaggccgct
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65 gttcaagctg cttcagcccc tcttattgaa cgctctattg aacaagcaac tggcattgta
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67 tacaacatca ctggtggacc ggacctcaca ttgcaggaag tcaacaccgt gtctgagatt
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69 gtaacaggtt tagctgaccc ctcagctaat atcatttttg gagcggtagt ggatgacaaa
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71 tatacaggtg aaatccatgt aacgattatt gccacggggt tctctcacag ttttcagaaa
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73 tcactagtgg acccaaacgt ttctaggtcg gagaggcagg acgccccgag taatgcactc
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75 gagaaacett ggaagcaace aacteecace teateaagat ttegteaagg cettaatage
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82 <212> TYPE: DNA
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90 agccaagtgc aatgccaacg ctttcgatgt ctatcttcag aatataaggg tcataatctt
                                                                         180
92 aaacttagaa gacgtagccg tgtctcagct tccaacagag aaaacggtag tttaaatggg
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94 cgtttccagg aatcactgag tcaagagaat gggtatccgg caccaactga agggactgat
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96 cctcacactt tctccacggc gatggactcc ttagctatta aagcagagga agcttacaat
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98 gacgtacagg attettttge caagagtagt aaacaaegga gettatetgg etgegettet
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102 gaactattga atgttgagtt ctgggccgtc aatactgaca aacaagcatt gaacaagtcg
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106 agaagtgcaa ccggtgagga agcagctaca gagtcattgg cggagctttc gatggcactt
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110 gctcctgtgg tggctcggtt ggcgaaggct atgggagcgt taacgattgg catagtaact
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116 gcacctgaca catctatgct ggaggctttc catcttgcag atgacgtctt gcggcaggga
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118 gtgcaaggaa tttcagacat catcacgata cccgggctag tcaacgtcga ctttgcggat
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120 gtgaaagcta tcatgtcaaa tgcagggagt gcaatgttgg gaatcggcgc tggttttggg
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122 aagaaccqtq ctgaggaggt ggcacqgtca gccatcatgt ctcctctact ccgctccgtc
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124 tcgagaccca tgggtattgt gtacaatgtg acaggtggga gcgacctaac tcttcacgag
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126 gtcaacatcg ctgccgaaat tgttcatgac atggctgatc caaacgcaaa tgttatcttt
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128 ggggcggtca ttgatgagag ctttaagggg atgatacgta tgactgtcat tgcaactgga
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130 tttagagagc ctggagagga gaaggtcgtt ggtagtgttc gaactgtaga cgatgatata
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132 ttctactggg aacagaataa gaataggtcc gaccttggca aagtgccgga cgttttgcga
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138 <211> LENGTH: 444
139 <212> TYPE: PRT
140 <213> ORGANISM: Physcomitrella patens
142 <400> SEQUENCE: 3
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147 Gly Ser Leu Cys Ser Thr Ser Pro Gln Ser Met His Pro Met Ser Ser
                20
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151 Val Ala Ala Lys Val Thr Arg Gln Cys Gly Cys Leu Arg Ala Gly Asn
           35
                                40
155 Lys Leu Asp Lys Asp Gln Phe Val Gly Asp Gly Lys Pro Leu Met His
156
                            55
159 Gln Gln Thr Arg Gly Trp Ser Gln Gly Arg Glu Arg Cys His Ala Gly
                                                                 80
160 65
                        70
                                            75
163 Arg Ser Val Val Met Ala Ser Met Ser Gly Ala Lys Ile Lys Val Ile
                                        90
167 Gly Val Gly Gly Gly Asn Asn Ala Val Asn Arg Met Ile Gly Ser
168
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171 Gly Ile Gln Gly Val Asp Phe Trp Ala Ile Asn Thr Asp Val Gln Ala 115 120 175 Leu Gln Lys Ser Gln Ala Glu His Arg Val Gln Ile Gly Glu Ala Leu 135 179 Thr Arg Gly Leu Gly Thr Gly Gly Lys Pro Phe Leu Gly Glu Gln Ala 183 Ala Glu Glu Ser Ile Glu Ile Ile Ala Gln Ala Val Val Asp Ala Asp 170 165 187 Leu Val Phe Ile Thr Ala Gly Met Gly Gly Thr Gly Ser Gly Ala 185 191 Ala Pro Val Val Ala Arg Val Ala Lys Glu Ala Gly Gln Leu Thr Val 195 200 195 Gly Val Val Thr Tyr Pro Phe Thr Phe Glu Gly Arg Arg Arg Ser Gln 196 210 215 199 Gln Ala Val Glu Ala Ile Glu Asn Leu Arg Lys Ser Val Asp Ser Leu 235 230 203 Ile Val Ile Pro Asn Asp Arg Leu Leu Asp Val Ser Gly Asp Lys Thr 245 250 207 Pro Leu Gln Glu Ala Phe Ser Leu Ala Asp Asp Val Leu Arg Gln Gly 208 265 211 Val Gln Gly Ile Ser Asp Ile Ile Thr Thr Pro Gly Leu Val Asn Val 212 275 280 215 Asp Phe Ala Asp Val Arg Ala Val Met Ser Asn Ser Gly Thr Ala Met 295 300 219 Leu Gly Val Gly Ser Ser Ser Gly Lys Asn Arg Ala Glu Glu Ala Ala 310 315 223 Val Gln Ala Ala Ser Ala Pro Leu Ile Glu Arg Ser Ile Glu Gln Ala 325 330 227 Thr Gly Ile Val Tyr Asn Ile Thr Gly Gly Pro Asp Leu Thr Leu Gln 340 345 231 Glu Val Asn Thr Val Ser Glu Ile Val Thr Gly Leu Ala Asp Pro Ser 360 235 Ala Asn Ile Ile Phe Gly Ala Val Val Asp Asp Lys Tyr Thr Gly Glu 375 239 Ile His Val Thr Ile Ile Ala Thr Gly Phe Ser His Ser Phe Gln Lys 390 395 243 Ser Leu Val Asp Pro Asn Val Ser Arg Ser Glu Arg Gln Asp Ala Pro 405 410 247 Ser Asn Ala Leu Glu Lys Pro Trp Lys Gln Pro Thr Pro Thr Ser Ser 420 425 251 Arg Phe Arg Gln Gly Leu Asn Ser Lys Gly Phe Leu 252 435 440 255 <210> SEQ ID NO: 4 256 <211> LENGTH: 490 257 <212> TYPE: PRT 258 <213> ORGANISM: Physcomitrella patens 260 <400> SEQUENCE: 4 261 Met Ile Thr Cys Arg Val Trp Val Gly Leu Gly Pro Val Ser Pro Ser 262 1

Input Set : A:\MBP-017xxSeqList.txt
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265 Leu Ile Leu Leu Pro Ser Lys Ser Asn Gly Glu Cys Val Leu Ser Ala 25 269 Arg Lys Ala Asp Trp Gly Leu Leu Ser Gln Val Gln Cys Gln Arg Phe 273 Arg Cys Leu Ser Ser Glu Tyr Lys Gly His Asn Leu Lys Leu Arg Arg 55 277 Arg Ser Arg Val Ser Ala Ser Asn Arg Glu Asn Gly Ser Leu Asn Gly 70 281 Arg Phe Gln Glu Ser Leu Ser Gln Glu Asn Gly Tyr Pro Ala Pro Thr 285 Glu Gly Thr Asp Pro His Thr Phe Ser Thr Ala Met Asp Ser Leu Ala 100 105 289 Ile Lys Ala Glu Glu Ala Tyr Asn Asp Val Gln Asp Ser Phe Ala Lys 115 120 293 Ser Ser Lys Gln Arg Ser Leu Ser Gly Cys Ala Ser Ile Lys Val Phe 135 297 Gly Val Gly Gly Gly Cys Asn Ala Val Asp Glu Met Val Arg Ser 150 155 301 Glu Leu Leu Asn Val Glu Phe Trp Ala Val Asn Thr Asp Lys Gln Ala 170 305 Leu Asn Lys Ser Leu Ala Pro Asn Lys Ile Gln Ile Gly Gln Asp Thr 185 309 Thr Ala Gly Arg Gly Ala Gly Gly Arg Ser Ala Thr Gly Glu Glu Ala 200 313 Ala Thr Glu Ser Leu Ala Glu Leu Ser Met Ala Leu Glu Gly Ala Asp 215 317 Leu Val Phe Ile Ala Ser Gly Met Gly Gly Thr Gly Ser Gly Ala 230 235 321 Ala Pro Val Val Ala Arg Leu Ala Lys Ala Met Gly Ala Leu Thr Ile 245 250 325 Gly Ile Val Thr Glu Pro Phe Thr Phe Glu Gly Phe Thr Arg Ala Arg 265 329 Gln Ala Arg Lys Ala Ile Glu Asp Met Arg His Ala Ala Asp Thr Val 275 280 333 Val Val Val Pro Asn Asp Arg Leu Leu Gln Thr Val Ala Pro Asp Thr 295 337 Ser Met Leu Glu Ala Phe His Leu Ala Asp Asp Val Leu Arg Gln Gly 310 315 341 Val Gln Gly Ile Ser Asp Ile Ile Thr Ile Pro Gly Leu Val Asn Val 325 330 345 Asp Phe Ala Asp Val Lys Ala Ile Met Ser Asn Ala Gly Ser Ala Met 345 349 Leu Gly Ile Ala Leu Val Leu Gly Lys Asn Arg Ala Glu Glu Val Ala 355 360 353 Arg Ser Ala Ile Met Ser Pro Leu Leu Arg Ser Val Ser Arg Pro Met 375 357 Gly Ile Val Tyr Asn Val Thr Gly Gly Ser Asp Leu Thr Leu His Glu 395 361 Val Asn Ile Ala Ala Glu Ile Val His Asp Met Ala Asp Pro Asn Ala

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362 405 410 365 Asn Val Ile Phe Gly Ala Val Ile Asp Glu Ser Phe Lys Gly Met Ile 425 369 Arg Met Thr Val Ile Ala Thr Gly Phe Arg Glu Pro Gly Glu Glu Lys 440 435 373 Val Val Gly Ser Val Arg Thr Val Asp Asp Asp Ile Phe Tyr Trp Glu 455 377 Gln Asn Lys Asn Arg Ser Asp Leu Gly Lys Val Pro Asp Val Leu Arg 470 475 381 Arg Lys Asp Arg Arg Gly Ser Gly Arg 485 490 385 <210> SEQ ID NO: 5 386 <211> LENGTH: 383 387 <212> TYPE: PRT 388 <213> ORGANISM: Physcomitrella patens 390 <400> SEQUENCE: 5 391 Met Asp Ser Leu Ala Ile Lys Ala Glu Glu Ala Tyr Asn Asp Val Gln 395 Asp Ser Phe Ala Lys Ser Ser Lys Gln Arg Ser Leu Ser Gly Cys Ala 399 Ser Ile Lys Val Phe Gly Val Gly Gly Gly Cys Asn Ala Val Asp 403 Glu Met Val Arg Ser Glu Leu Leu Asn Val Glu Phe Trp Ala Val Asn 55 407 Thr Asp Lys Gln Ala Leu Asn Lys Ser Leu Ala Pro Asn Lys Ile Gln 411 Ile Gly Gln Asp Thr Thr Ala Gly Arg Gly Ala Gly Gly Arg Ser Ala 415 Thr Gly Glu Glu Ala Ala Thr Glu Ser Leu Ala Glu Leu Ser Met Ala 100 105 419 Leu Glu Gly Ala Asp Leu Val Phe Ile Ala Ser Gly Met Gly Gly Gly 423 Thr Gly Ser Gly Ala Ala Pro Val Val Ala Arg Leu Ala Lys Ala Met 135 427 Gly Ala Leu Thr Ile Gly Ile Val Thr Glu Pro Phe Thr Phe Glu Gly 155 431 Phe Thr Arg Ala Arg Gln Ala Arg Lys Ala Ile Glu Asp Met Arg His 165 170 435 Ala Ala Asp Thr Val Val Val Pro Asn Asp Arg Leu Leu Gln Thr 185 439 Val Ala Pro Asp Thr Ser Met Leu Glu Ala Phe His Leu Ala Asp Asp 195 200 443 Val Leu Arg Gln Gly Val Gln Gly Ile Ser Asp Ile Ile Thr Ile Pro 215 210 220 447 Gly Leu Val Asn Val Asp Phe Ala Asp Val Lys Ala Ile Met Ser Asn 230 235 451 Ala Gly Ser Ala Met Leu Gly Ile Ala Leu Val Leu Gly Lys Asn Arg 455 Ala Glu Glu Val Ala Arg Ser Ala Ile Met Ser Pro Leu Leu Arg Ser

VERIFICATION SUMMARYDATE: 02/22/2006PATENT APPLICATION: US/10/538,530TIME: 12:11:08

Input Set : A:\MBP-017xxSeqList.txt
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